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OM nucleic - nucleic search, using sw model

Run on: August 19, 2003, 12:03:50 ; Search time 576 Seconds

(Without alignments)  
8635.202 Million cell updates/sec

Title: US-09-494-297-1

Perfect score: 2274

Sequence: 1 atgaaaaaaccaaggttc.....ggataagaacaatgactag 2274

Scoring table: IDENTITY\_NUC

Gapop 10.0 , Gapept 1.0

Searched: 1504479 seqs, 1118970152 residues

Total number of hits satisfying chosen parameters: 3008958

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Listing first 45 summaries

Database : Published\_Applications\_NA:\*

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16: /cgn2_6/ptodata/2/pubpna/US60_PUBCOMB.seq:*
17: /cgn2_6/ptodata/2/pubpna/US60_PUBCOMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	53	2.3	335913	11	US-09-754-853A-2
2	53	2.3	335913	11	US-09-754-853A-3
3	52.4	2.3	6106	12	US-10-311-455-1445
4	52	2.3	7195	12	US-10-240-453-40
5	52	2.3	7195	14	US-10-239-676-30
6	51.2	2.3	4985	14	US-10-094-240-10
7	50.6	2.2	6815	12	US-10-311-455-644
8	50.6	2.2	6815	12	US-10-240-453-50
9	50.6	2.2	6815	14	US-10-239-676-50
10	50.6	2.2	7819	12	US-10-311-455-1926
11	50.6	2.2	7819	12	US-10-240-485-160
12	50.2	2.2	19087	12	US-10-311-455-766
13	50	2.2	2000	10	US-09-938-842A-2735
14	50	2.2	6381	12	US-10-311-455-939
15	50	2.2	6381	12	US-10-240-485-71
16	50	2.2	640681	10	US-09-790-988-1

c 17	49.6	2.2	12705	12	US-10-311-455-121	Sequence 121, App
c 18	49	2.2	16724	12	US-10-311-455-1063	Sequence 1063, Ap
c 19	49	2.2	16724	12	US-10-240-485-89	Sequence 89, Appl
c 20	48.2	2.1	5539	12	US-10-240-485-180	Sequence 180, Appl
c 21	48.2	2.1	6277	12	US-10-311-455-2013	Sequence 2013, Ap
c 22	48.2	2.1	11944	12	US-10-311-455-2160	Sequence 2160, Ap
c 23	48	2.1	9728	12	US-10-311-455-1876	Sequence 1876, Ap
c 24	47.8	2.1	6306	12	US-10-204-708-50	Sequence 50, Appl
c 25	47.8	2.1	6712	12	US-10-311-455-664	Sequence 664, App
c 26	47.4	2.1	5897	8	US-08-781-986A-320	Sequence 320, App
c 27	47.2	2.1	414	10	US-09-960-352-6528	Sequence 6528, Ap
c 28	47	2.1	625	13	US-10-027-632-250075	Sequence 250075, Sequence 1642, Ap
c 29	47	2.1	5347	12	US-10-311-455-1642	Sequence 2143, Ap
c 30	47	2.1	6103	12	US-10-311-455-2143	Sequence 2143, Ap
c 31	47	2.1	11260	14	US-10-239-676-19	Sequence 19, Appl
c 32	47	2.1	11260	14	US-10-311-455-1271	Sequence 1271, Ap
c 33	47	2.1	12025	12	US-10-311-455-1919	Sequence 1919, Ap
c 34	47	2.1	16811	12	US-10-240-453-281	Sequence 281, App
c 35	46.8	2.1	9504	12	US-10-311-455-588	Sequence 588, App
c 36	46.8	2.1	13584	12	US-10-311-455-2128	Sequence 2128, App
c 37	46.8	2.1	15548	12	US-10-311-455-227	Sequence 227, App
c 38	46.8	2.1	18154	12	US-09-754-853A-3	Sequence 3, Appl
c 39	46.8	2.1	335913	11	US-09-754-853A-2	Sequence 67, Appl
c 40	46.8	2.1	335913	11	US-10-311-455-67	Sequence 5, Appl
c 41	46.6	2.0	7928	12	US-10-311-455-336	Sequence 336, App
c 42	46.6	2.0	10640	12	US-10-240-485-30	Sequence 30, Appl
c 43	46.6	2.0	15674	12	US-10-311-455-76212	Sequence 76212, A
c 44	46.6	2.0	15674	12	US-10-240-485-30	
c 45	46.6	2.0	2140405	13	US-10-027-632-76212	

## ALIGNMENTS

```
RESULT 1
US-09-754-853A-2
: Sequence 2, Application US/09754853A
: Publication No. US20030005491A1
GENERAL INFORMATION:
: APPLICANT: Hauge, Brian M.
: APPLICANT: Parnell, Laurence D.
: APPLICANT: Parsons, Jeremy D.
: APPLICANT: Wang, Ming Li
: TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
: TITLE OF INVENTION: Soybean Cyst Nematode Resistance
: FILE REFERENCE: 38-10(13810)B
: CURRENT APPLICATION NUMBER: US/09/754, 853A
: PRIOR FILING DATE: 2001-01-05
: PRIOR APPLICATION NUMBER: US 60/174, 880
: PRIOR FILING DATE: 2000-01-07
: NUMBER OF SEQ ID NOS: 1119
SEQ ID NO 2
: LENGTH: 335913
: TYPE: DNA
: ORGANISM: Glycine max
: FEATURE:
: NAME/KEY: CDS
: LOCATION: (45163)..(45314)..(45450)..(45509)..(46941)..(48763)..(48975)..(49573)
: OTHER INFORMATION: Clone ID: 240017_region_G3
US-09-754-853A-2
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Query Match Best Local Similarity 2.3% Score 53: DB 11: Length 335913:

Matches 185: Conservative 0: Mismatches 220: Indels 0: Gaps 0:

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OY 1712 ACAATAAATATCAATCTTATTCGACTGATGATCCAGAGATTAGTATATTA 1771
||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 99367 ACAAAATATTAATATTAATATTAATCTCAATTTAAATATACAAATATTTAAATTA 99426
||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
OY 1772 TTTCGTATGAGATTAATTAATTAATCTGTAATCTGTAATTAATTAATGAGAAA 1831
||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 99427 AACATTTATTAATAAATAATATTAATTAATTAATTAATTAATTAATTAATTA 99486
||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
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Oy	1832	CGGAGCATGGTTTAGTGGTGACAGAACTTAATTTCCATTGTGAATTCGAAATTTAAAAA	1891
Dd	99487	AATCATTTAAAAAATACATACCAAAATRTTAAATAAACATPATAGAAGTBAATTTAAAAA	99546
Oy	1892	ATAATTAAGCAAGATTGCCTTCTCCAACCTGTAAAAACAGATAAAACAACCTCGAATTTA	1951
Dd	99547	TATTTATTTATTAATTAATTAATAAATTTATTTACACTTTTAAAAAATTTGAAAATA	99606
Oy	1952	AAGATGGTAAAGCACACATTAAATTTAAACATGGGAAGTTTAACACTCAAGCTTTC	2011
Dd	99607	AAAACAAAAAAGAAAATATCAAGTBAAATAAAAATCTBAAAACAAAAACTTTAAACATGTA	99666
Oy	2012	CAGAAGTTATTCCTTACCTTGTCAAGAAACAGATTCGTGAGGCTTTAAGTTTAAAGTTA	2071.
Dd	99667	AAAAAATGACTTTAAATCAATAGAAAAAATAAACTTACAACTTCAAACTGGTAAGTTCA	99726
Oy	2072	ATAGCCAAAGAATRGCAAAATGCTACAGCTTCCAAAAACAGAAATTA	2116
Dd	99727	AAGAAAAAACCCTTGACGACTTTTAAAAATGTAAATTTAAAAATAAAAA	99771

## RESULT 2

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US-09-754-853A-3
Sequence 3, Application US/09754853A
Publication No. US20030005491A1
GENERAL INFORMATION:
APPLICANT: Hauge, Brian M.
APPLICANT: Parnell, Laurence D.
APPLICANT: Parsons, Jeremy D.
APPLICANT: Wang, Ming Li
TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
FILE REFERENCE: 38-10(15810)B
CURRENT APPLICATION NUMBER: US/09/754, 853A
CURRENT FILING DATE: 2001-01-05
PRIORITY APPLICATION NUMBER: US 60/174, 880
PRIORITY FILING DATE: 2000-01-07
NUMBER OF SEQ ID NOS: 1119
SEQ ID NO 3
LENGTH: 335913
TYPE: DNA
ORGANISM: Glycine max
FEATURE:
NAME/KEY: CDS
LOCATION: (46798)..(48763),(48975)..(49573)
OTHER INFORMATION: Clone ID: 240017_region_G3
US-09-754-853A-3

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Query Match	Score	DB	Length
2.38;	53;	11;	335913;

Best Local Similarity 45.7%; Pred. No. 4.9;  
Matches 185; Conservative 0; Mismatches 220; Indels 0; Gaps 0;

QY	1712	ACAAATTAATATCATCTCTTTATTTGGAACTCGTGGCAATCCGGAAGATTATTAATTA	1771
Db	99367	ACAAAATATTTAAATATATATAAATTAATCTCAATTTTAAATATACAAAATATATTTAAATAA	9422
QY	1772	TTGCTATGGAAGATAAAAAGAACTTATACCTGTAAGTCAATTAATTTAAACATTTGAGAAAA	1831
Db	99427	AACATTTATTTAAAAATATATATAAATTTTAAAAATATAATATATATTAATTAATTA	9448
QY	1832	CGGTGAAGTGGTTTGCTGGTGAGCAAGTAAGTTCCATTTTGAATTTGAATTAATTA	1891
Db	99487	AATCATTTAAAAAAATACATACAAATATTTAAATAAAACTAATTAAGAACTAAATTAATAAA	9954
QY	1892	ATAATTAAGCAAGATTGCTTTCTCAACGTAAAAAGATAAAAACAACCTCGAATTTA	1951
Db	99547	TATTTATTTATATATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTA	9960
QY	1952	AAGTGGTAAAGCAACCATTAATTTAAACAATGGGGAAAGTTTAAACACTCAAGGTTAC	2011
Db	99607	AAAACAAAAGAAATATACAGTAAATTAATTAATTTCTTAACAAACAAACCTTTAAACATGAA	9966
QY	2012	CAGAAGTTATTTCTACCTTGTCAAAGAAACAGATTTCTGAAGGCTATTAAGGTTAAAGTTA	2071

Db 99667 AAAAAAAAAATGACTTTAAAGTCATAGAAAAAAAAAAAAAAGCTTACACTTCCTCAAGTGGTAAGTTCA 99726  
 Oy 2072 ATAGGCAAGAGTAGCAAAATGCTACAGTTTCAAAAAACAGAAATA 2116  
 Db 99727 AAGAAAAAAGCTGTAGAGCTTTAAAAATGTAAATTTAAAAATAAAAA 99771

### RESULT 3

US-10-311-455/445/c  
Sequence 1445, Application US/10311455  
Publication No. US20030143606A1  
GENERAL INFORMATION:  
APPLICANT: OLEK, Alexander  
APPLICANT: PIPENBROCK, Christian  
APPLICANT: BERLIN, Kurt  
TITLE OF INVENTION: Diagnosis of Diseases Associated with the Immune System by Det  
TITLE OF INVENTION: Cytosine methylation  
FILE REFERENCE: 5013.1014  
CURRENT APPLICATION NUMBER: US/10/311.455  
CURRENT FILING DATE: 2002-12-16  
PRIOR APPLICATION NUMBER: PCT/EP01/07537  
PRIOR FILING DATE: 2001-07-02  
PRIOR APPLICATION NUMBER: DE 10032529.7  
PRIOR FILING DATE: 2000-06-30  
PRIOR APPLICATION NUMBER: DE 10043826.1  
PRIOR FILING DATE: 2000-09-01  
NUMBER OF SEQ ID NOS: 2424  
SEQ ID NO 1445  
LENGTH: 6106  
TYPE: DNA  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)  
US-10-311-455-1445

Query Match 2.38; Score 52.4; DB 12; Length 6106;

Best Local Similarity 46.7%; Pred: No. 1.1;  
Matches 271; Conservative 0; Mismatches 301; Indels 8; Gaps 3

QY	1544	TATTTTATTTTCACAGTAGTCGCGTAATTTAGTAAAGCAATTAAGCAATTCATGGTT	1603
Db	2939	TATATATATTTATATATATTTATTTATATATATATATATATATATATATATATATAT	2880
QY	1604	TTGGAGACATCAATGATAGTACTTTAGCAGTTGCTGTAATAATCCTTGTAGATAATCGTC	1663
Db	2879	ATATCAT	2820
QY	1664	ATAGTATATCCCGCCAGCGTACATCGACCGTGGTTTCTTATATCCGAATATACATAATTC	1723
Db	2819	TTAT	2761
QY	1724	AATCTCTTATTTGGACTCAGTGGCATCGACAGATTTAGTATATATATATTTGATGGA	1783
Db	2760	ATTATATTTAT	2701
QY	1784	ATATAAAAAAGAGTTATACCTGTACCTATATTTTACATTTAGAGAAAAACGCTGACTG	1843
Db	2700	ATATTTAAAAAACTTAAATATATTAACATTTATATATACATTTT-----AATPATTTTCTTTAA	2646
QY	1844	TAGCTGGTGCAGACATTAAGATTTCCATTTTGAATTTAAAAATATATATAGCAAG	1903
Db	2645	CTCTTAAAAAACCAAAAAACCATTTAAATATAAAAACATTTCTCATATAAAAAAACAAAT	2586
QY	1904	AATTCCTTTCTCAACGTGTATTAACACATATAAACCAAACTCGAATTTTAAAGATGCTAAG	1963
Db	2585	AACTAATTTATTTATTTACTTTATTCACATCACTAATTTATCTTATTTAAAAATAAACTAT	2526
QY	1964	CAACCATTTATTTAAACATATGGGGAAGTTTAACTGTCGAAGTTTTCGCCGAAGGTTAT	2023
Db	2525	CAAAAACAAAAAATCAAAAAAACCAATCAAAAAACATATATTCACACATTTCAAAAAATA	2466
QY	2024	CTTACCTTGTCAAAAGAAACGATTTCTGAAGGCTTTAAGGTTTAAAGTTAATATGCCAAG	2083

Db 2465 ATAACTTAATCAAAAAAAGCTATTAACCTAACCAATTA--CTAAAAAAGCTAAAAA 2408  
QY 2084 TAGCAATGCTACAGCTTTCAAAAACAGAAATACAGTGA 2123  
Db 2407 TAAAAAATAAAAATTTTAAACGATCTATATAAATA 2368

## RESULT 4

US-10-240-453-40/C  
; Sequence 40, Application US/10240453  
; Publication No. US20030148326A1  
; GENERAL INFORMATION:  
; APPLICANT: OLEK, Alexander  
; APPLICANT: PIEPENBROCK, Christian  
; APPLICANT: BERLIN, Kurt  
; TITLE OF INVENTION: Diagnosis of Diseases Associated with DNA  
; TITLE OF INVENTION: Transcription  
; TITLE OF INVENTION: by Means of Assessing the Methylation Status of Genes Associated  
; FILE REFERENCE: 5013.1009  
; CURRENT APPLICATION NUMBER: US/10/240,453  
; CURRENT FILING DATE: 2002-10-02  
; PRIOR APPLICATION NUMBER: PCT/EP01/03973  
; PRIOR FILING DATE: 2001-04-06  
; PRIOR APPLICATION NUMBER: DE 10019058.8  
; PRIOR FILING DATE: 2000-04-06  
; PRIOR APPLICATION NUMBER: DE 10019173.8  
; PRIOR FILING DATE: 2000-04-07  
; PRIOR APPLICATION NUMBER: DE 10032529.7  
; PRIOR FILING DATE: 2000-06-30  
; PRIOR APPLICATION NUMBER: DE 10043826.1  
; PRIOR FILING DATE: 2000-09-01  
; NUMBER OF SEQ ID NOS: 350  
; SEQ ID NO 40  
; LENGTH: 7195  
; TYPE: DNA  
; ORGANISM: Artificial Sequence  
; FEATURE:  
; OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)  
US-10-240-453-40

Query Match 2.3%; Score 52; DB 12; Length 7195;  
Best Local Similarity 52.3%; Pred. No. 1.4;  
Matches 139; Conservative 0; Mismatches 125; Indels 2; Gaps 1;

QY 1717 AATATCACTCTTATGGACCTGAGTCAGATCCAGAGATTAGTGAATTAATTCGT 1776  
Db 1246 AAACATTTAAATTTTAAATATCTCAATATATTCACATTAACATTAATTAATCTA 1187  
QY 1777 ATGAGATTAATAAAGAGTTATACCTTAATCTAATTTAACAATTGAGAAAAACGGTG 1836  
Db 1186 AACACAAAAAATAAATAATATCTTAATTAATTAACAATAAATAAATAAATAA 1127  
QY 1837 ACTGTTAGCTGTGACAGAACTTAAGATTTCATTTGAATTTGAATTAATAAAT 1896  
Db 1126 ATAAATATAAACAATTAATAAATAAAGACATCATCTTAATTTAATAATAAAT 1067  
QY 1897 AAGCAAGATTCCTTCTCAACCTGTATAACAGATAAACAACCTCGAATTAAGAT 1956  
Db 1066 AAAATATATATTAATTAACATAACCTTCCAAAATATATTTAAACACTTCTTAA--AT 1009  
QY 1957 GGTAAAGCAACCATTAATTTAAACA 1982  
Db 1008 CCTAAAAAATCTTAATTTAAATA 983

RESULT 5  
US-10-239-676-30/C  
; Sequence 30, Application US/10239676  
; Publication No. US20030082609A1  
; GENERAL INFORMATION:  
; APPLICANT: OLEK, Alexander

APPLICANT: PIEPENBROCK, Christian  
APPLICANT: BERLIN, Kurt  
TITLE OF INVENTION: Diagnosis of Diseases Associated with Gene Regulation  
FILE REFERENCE: 5013.1003  
CURRENT APPLICATION NUMBER: US/10/239,676  
CURRENT FILING DATE: 2002-09-24  
PRIOR APPLICATION NUMBER: PCT/EP01/03968  
DE 10019058.8  
DE 10019173.8  
DE 10032529.7  
DE 10043826.1  
PRIOR FILING DATE: 2001-04-06  
PRIOR FILING DATE: 2000-04-06  
PRIOR FILING DATE: 2000-04-07  
PRIOR FILING DATE: 2000-06-30  
PRIOR FILING DATE: 2000-09-01  
NUMBER OF SEQ ID NOS: 228  
SEQ ID NO 30  
LENGTH: 7195  
TYPE: DNA  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)  
US-10-239-676-30

Query Match 2.3%; Score 52; DB 14; Length 7195;  
Best Local Similarity 52.3%; Pred. No. 1.4;  
Matches 139; Conservative 0; Mismatches 125; Indels 2; Gaps 1;

QY 1717 AATATCACTCTTATGGACCTGAGTCAGATCCAGAGATTAGTGAATTAATTCGT 1776  
Db 1246 AAACATTTAAATTTTAAATATCTCAATATATTCACATTAACATTAATTAATCTA 1187  
QY 1777 ATGAGATTAATAAAGAGTTATACCTTAATCTAATTTAACAATTGAGAAAAACGGTG 1836  
Db 1186 AACACAAAAAATAAATAATATCTTAATTAATTAACAATAAATAAATAAATAA 1127  
QY 1837 ACTGTTAGCTGTGACAGAACTTAAGATTTCATTTGAATTTGAATTAATAAAT 1896  
Db 1126 ATAAATATAAACAATTAATAAATAAAGACATCATCTTAATTTAATAATAAAT 1067  
QY 1897 AAGCAAGATTCCTTCTCAACCTGTATAACAGATAAACAACCTCGAATTTAAGAT 1956  
Db 1066 AAAATATATATTAATTAACATAACCTTCCAAAATATATTTAAACACTTCTTAA--AT 1009  
QY 1957 GGTAAAGCAACCATTAATTTAAACA 1982  
Db 1008 CCTAAAAAATCTTAATTTAAATA 983

RESULT 6  
US-10-094-240-10/C  
; Sequence 10, Application US/10094240  
; Publication No. US20030082637A1  
; GENERAL INFORMATION:  
; APPLICANT: ZWIBEL, LAURENCE J.  
; TITLE OF INVENTION: ARRESTIN GENE, POLYPEPTIDE, AND METHODS OF USE THEREOF  
; FILE REFERENCE: N8289  
; CURRENT APPLICATION NUMBER: US/10/094,240  
; CURRENT FILING DATE: 2001-03-08  
; PRIOR APPLICATION NUMBER: 10/056,405  
; PRIOR FILING DATE: 2002-01-24  
; PRIOR APPLICATION NUMBER: 60/264,649  
; PRIOR FILING DATE: 2001-01-26  
; NUMBER OF SEQ ID NOS: 27  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 10  
; LENGTH: 4985  
; TYPE: DNA  
; ORGANISM: Anopheles gambiae  
US-10-094-240-10

Query Match 2.3%; Score 51.2; DB 14; Length 4985;

Best Local Similarity 43.98; Pred. No. 1.7;  
Matches 266; Conservative 0; Mismatches 338; Indels 2; Gaps 1;

OY	1324	AAAACATGACTCCAGCTTTTACCACAACGGAGCACTAAATAATCAGCTCATTTGCAGGTGCT	1383
Dd	2924	ATTATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	2865
OY	1384	GACCTCTTTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	1443
Dd	2864	AATAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	2805
OY	1444	ATCAAAAAAGTAATTGGAGAGGGTTACAGGAAAAGACAGCTATTGAGTATAGTGCT	1503
Dd	2804	AAGAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	2745
OY	1504	CTAACGAGACACATTTGCCGCGCTACTCAGTTAGCAATATATTTATTCAGTAGTAGT	1563
Dd	2744	ATTATTTTATTGATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	2685
OY	1564	GCTGAATTTAGATAAGATTAACCTAAAGACTATCATGCTGTTTGAGACATGATGATAGT	1623
Dd	2684	AATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	2627
OY	1624	ACTTTAGCAGTGGCTAAATCCTTGTAATTAACCTCAAGTAACTAATCCGCCACAGCTA	1683
Dd	2626	AATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	2567
OY	1684	ACTGACCTTGATTTCTTTATTCGCAATACAACTAAATATCAATCTCTTATGGAACCTCAG	1743
Dd	2566	AATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	2507
OY	1744	TGGCATCCAGAGATTAGTATTTTCGTATTTGATGGAAGATTAAGAAAAGAGATTATACCT	1803
Dd	2506	AATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	2447
OY	1804	GTACGCTATTAATTAACATTGGAGAAAAAGSGTGCTGGTTAGCTGGGACAGAACTAAA	1863
Dd	2446	AATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	2387
OY	1864	GATTTCCATTTTGAATTTGAATTTAAAAAATTAATTAAGCAGAAATGCTTTCCAAACTGTT	1923
Dd	2386	AATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT	2327
OY	1924	AAAAACA 1929	
Dd	2326	ACTAAA 2321	

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RESULT 7
US-10-311-455-644/C
Sequence 644, Application US/10311455
Publication No. US20030143606A1
GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
APPLICANT: PIRENBERG, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with the Immune System by Determining the Cytosine Methylation of Cytosine Methylation
FILE REFERENCE: 5013.1014
CURRENT APPLICATION NUMBER: US/10/311.455
CURRENT FILING DATE: 2002-12-16
PRIORITY APPLICATION NUMBER: PCT/EP01/07537
PRIORITY FILING DATE: 2001-07-02
PRIORITY APPLICATION NUMBER: DE 10032529.7
PRIORITY FILING DATE: 2000-06-30
PRIORITY APPLICATION NUMBER: DE 10043826.1
PRIORITY FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 2424
SEQ ID NO 644
LENGTH: 6815
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:

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; OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)  
US-10-311-455-644

[illegible]

```

RESULT 8
US-10-240-453-50/c
; Sequence 50, Application US/10240453
; Publication No. US20030148326A1
; GENERAL INFORMATION:
; APPLICANT: OLEK, Alexander
; APPLICANT: PIEPENBROCK, Christian
; APPLICANT: BERLIN, Kurt
; TITLE OF INVENTION: Diagnosis of Diseases Associated with DNA
; TITLE OF INVENTION: Transcription
; TITLE OF INVENTION: by Means of Assessing the Methylation Status of Genes Associated
; TITLE OF INVENTION: with DNA Transcription
; FILE REFERENCE: 5013.1009
; CURRENT APPLICATION NUMBER: US/10/240.453
; CURRENT FILING DATE: 2002-10-02
; PRIOR APPLICATION NUMBER: PCT/EP01/03973
; PRIOR FILING DATE: 2001-04-06
; PRIOR APPLICATION NUMBER: DE 10019058.8
; PRIOR FILING DATE: 2000-04-06
; PRIOR APPLICATION NUMBER: DE 10019173.8
; PRIOR FILING DATE: 2000-04-07
; PRIOR APPLICATION NUMBER: DE 10032529.7
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: DE 10043826.1
; PRIOR FILING DATE: 2000-09-01
; NUMBER OF SEQ ID NOS: 350
; SEQ ID NO 50
; LENGTH: 6815
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
US-10-240-453-50

```

	Query Match	2.2%	Score 50.6;	DB 12;	Length 6815;
	Best Local Similarity	48.8%;	Pred. No. 2.6;		
	Matches 137;	Conservative 0;	Mismatches 144;	Indels 0;	Gaps 0;
OY	1871	ATTTGAATTGAAATTAATAAAGCAGACGATTCCTTCCAAACTGTAAAAACG	1930		
Dd	5735	AATTTAAATTAATTAATCGAAAAAAAACCAAAATTAATCCTTTAAAAAACGGTTAAAAACA	5676		
OY	1931	ATAAACCAACCTTGAAATTTAAAGCATGTTAAAGCAACCATTAATTTAAAAACATGGGGAAA	1990		
Dd	5675	ATAAAATTAATTAATCTCAAAAAAAATTAACGGAATTCAGATTAATAAAAAAAAAAATTTATAAA	5616		

QY	1991	GTTTAAACCTCCAAAGGTTTACCCAGAGSAGTATTTCTACCTGTGTCAAGGAACAGATCTG	2050
Db	5615	ACTTAAAAACCTTAAAAATTAACCTCTTAATTAAATATATAAAAACAAAAACAATTAACCT	5556
QY	2051	AAGCCTATTAAGGTTAAAGTTAATAGCCACAGAGTACGAATGCTACAGTTTCAAAAACAG	2110
Db	5555	CGAAAAAACCAATAAATTTTAACTCTAAATTAACATAAAAAACAATAATTAACACACAAA	5496
QY	2111	GAATTAACAGTGTAGACACTTGTCTTTGTGAATAATAATAA	2151
Db	5495	CACGAAAAAATTAAAAAAACACCATTATTTTAAAAATATATA	5455

```

: RESULT 9
: US-10-239-676-50/c
: Sequence 50, Application US/10239676
: Publication No. US20030082609A1
: GENERAL INFORMATION:
: APPLICANT: OLEK, Alexander
: APPLICANT: PIEPENBROCK, Christian
: APPLICANT: BERLIN, Kurt
: TITLE OF INVENTION: Diagnosis of Diseases Associated with Gene Regulation
: FILE REFERENCE: 5013.1003
: CURRENT APPLICATION NUMBER: US/10/239,676
: CURRENT FILING DATE: 2002-09-24
: PRIORITY APPLICATION NUMBER: PCT/EP01/03968
: DE 10019058.8
: DE 10019173.8
: DE 10032529.7
: DE 10043826.1
: PRIORITY FILING DATE: 2001-04-06
: 2000-04-06
: 2000-04-07
: 2000-06-30
: 2000-09-01
: NUMBER OF SEQ ID NOS: 228
: SEQ ID NO 50
: LENGTH: 6815
: TYPE: DNA
: ORGANISM: Artificial Sequence
: FEATURE:
: OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
: US-10-239-676-50

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Query Match	2.2%	Score 50.6	DB 14	Length 6815
Best Local Similarity	48.8%	Pred. No. 2.6		
Matches 137	Conservative	0	Mismatches 144	Indels 0
				Gaps 0

QY	1871	ATTTTGGAATTSGATTTAAAAAATTAATTAAGCAGAATGGTTTCGCAACGTTGTAACAG	1930
Db	5735	AATTTAAATTAATAATCCGAAAAAANAACAAAATTAATCTTTAAAAACGGTTAAACAA	5676
QY	1931	ATAAAAAACAACCTCGAATTTAAGATGTAAGCAACCATTAATTTAAAAACATGGGAAA	1990
Db	5675	ATAAAAAATAAATCTCAAAAAAATAAACCGAAATACAAATAAAAAANAATAATATAA	5616
QY	1991	GTTTAACACTCAAGTTTACCGAGAGATATCTTACCTTGTCACAAGAAACAGATCTG	2050
Db	5615	ACTAAAAAACTTAAAAATTACCTCTTAATTAATATATAAAAACAAAAAATCAACCTT	5556
QY	2051	AAGCCTTAAGGTTAAAGTTAATAGCCACAGAGTACAAATGCTACAGTTTCAAAAAACAG	2110
Db	5555	CGAAAAAANAACAATTAATTTTAATCTAAATTAAGTAAAAAANAACATTAATACTAAMCAGAA	5496
QY	2111	GAATACAAAGTGAAGACACTTGCTTTGAAAAATATAAA	2151
Db	5495	CACGAAAAAATTTAAAAAACAACCATATTTTAAAAATATATA	5455

RESULT 10  
US-10-311-455-1926/c  
; Sequence 1926, Application US/10311455

Publication No. US20030143606A1  
GENERAL INFORMATION:  
APPLICANT: OLEK, Alexander  
APPLICANT: PIPENBROCK, Christian  
APPLICANT: BERLIN, Kurt  
TITLE OF INVENTION: Diagnosis of Diseases Associated with the Immune System by Det  
TITLE OF INVENTION: Cytosine methylation  
FILE REFERENCE: 5013.1014  
CURRENT APPLICATION NUMBER: US/10/311.455  
CURRENT FILING DATE: 2002-12-16  
PRIOR APPLICATION NUMBER: PCT/EP01/07537  
PRIOR FILING DATE: 2001-07-02  
PRIOR APPLICATION NUMBER: DE 10032529.7  
PRIOR FILING DATE: 2000-06-30  
PRIOR APPLICATION NUMBER: DE 10043826.1  
PRIOR FILING DATE: 2000-09-01  
NUMBER OF SEQ ID NOS: 2424  
SEQ ID NO 1926  
LENGTH: 7819  
TYPE: DNA  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)  
US-10-311-455-1926

Query Match 2.2%; Score 50.6; DB 12; Length 7819;  
Best Local Similarity 47.1%; Pred. No. 2.8;  
Matches 155; Conservative 0; Mismatches 174; Indels 0; Gaps 0;

QY	1638	TAAATTCCTTGTAATATACGTCAAGATAGTATATCTCCACAGCTACTGACCTTGATT	1693
Db	846	TAAATCCCAAAATTTAAATTTCAAAATATAAACCACTAGCCATCTCACTCACTT	787
QY	1698	CTTATTCGCAATPAACAATAATATCATCTCTATTGGAACCTAGTGGCATCCAGAAGA	1753
Db	786	TATTACTATATTTTAAATTCCTTTTACCTATATCTTTCAAACCAATTTTAACTATATCT	727
QY	1758	TTTATGTTGATTTATTTGCTATGGAAGATAAAAAAGATTTATACCTGATCATATATTT	1817
Db	726	ATCCCTAATATATATCTTATATATATACCAACCAAAAAAACTCCAGCTTACCCGAAACT	667
QY	1818	AACATGTGAAAAACGGTACTGGTTTAGCTGGTGACAGAACTAAAGATTTTCATTTTGA	1877
Db	666	AACATCAAAAACACTCAATATATCTAATACCACTCAAAACCTATTAACCAACAAAAAAT	607
QY	1878	AATTGAATTAATAAATATATTAAGCAGAAATTCCTTTCAAACTGTTAAACAGATTAAC	1933
Db	606	TATTAATTTAAAAATAAATAAATAATTAATTAATAAAAAAACCAACAAAAAATAAAAAA	547
QY	1938	AAACCTGAATTTAAAGATGTTAAAGCAA	1966
Db	546	AAAAAAAAAAAAAATCTTAAAAAATAA	518

US-RESULT 11  
US-10-240-485-160/C  
Sequence 160, Application US/10240485  
Publication No. US20030148327A1  
GENERAL INFORMATION:  
APPLICANT: OLER, Alexander  
APPLICANT: PIEPENBROCK, Christian  
APPLICANT: BERLIN, Kurt  
TITLE OF INVENTION: Diagnosis of Diseases  
TITLE OF INVENTION: Metastasis  
FILE REFERENCE: 5013.1007  
CURRENT APPLICATION NUMBER: US/10/240,485  
CURRENT FILING DATE: 2002-10-02  
PRIOR APPLICATION NUMBER: PCT/EP01/03970  
PRIOR FILING DATE: 2001-04-06  
PRIOR APPLICATION NUMBER: DE 10019058.8  
PRIOR FILING DATE: 2000-04-06  
PRIOR APPLICATION NUMBER: DE 10019173.8  
PRIOR FILING DATE: 2000-04-07



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? PRIOR APPLICATION NUMBER: DE 10032529.7
? PRIOR FILING DATE: 2000-06-30
? PRIOR APPLICATION NUMBER: DE 10043826.1
? PRIOR FILING DATE: 2000-09-01
? NUMBER OF SEQ ID NOS: 202
? SEQ ID NO 71
? LENGTH: 6381
? TYPE: DNA
? ORGANISM: Artificial Sequence
? FEATURE:
? OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
US-10-240-485-71

Query Match      2.2%; Score 50; DB 12; Length 6381;
Best Local Similarity 48.3%; Pred. No. 3.4;
Matches 140; Conservative 0; Mismatches 150; Indels 0; Gaps 0;

OY 1708 AATACAAATAAATATTCATCTCTTATTGGAACTGAGTCGACGACGAAGATTCTAGTAT 1767
    ||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1154 AATTACAAAAACAATTAACAAAAAATTTATCCCCCAAAATACCTAATTAACATTTCACA 1095

OY 1768 AATTATTCGTATGACAGATAAAAAAGAAGTTATFACCTGTAACATCAATTTAACATTGAGA 1822
    ||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1094 AATTACTCTTCATATAAAAAAATAATATATATACAAACAAATTACTATATCAAAAAAACA 1033

OY 1828 AAAACGGTACGTGTTAGCTGCGTGACAGACACTAAAGTTTCCATTTTGAAGATTGAATTA 1887
    ||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1034 AAAAAATTAATAATACAAATTAATTAATACACAACTCCAAACCAAAATTTATTTACATATAATA 975

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QY      1888  AAAAATATATAGCAGAGATGCTTGTCTGCACATCTGTAAAACAGATATAAACAAACCTGGAA 1947
          ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db      974  ATATTATTCATATATATATATATTTTAAAAAATTTTAAAAATTCAAAATCTAAATACCTBA 915
          ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY      1948  TTTTAAAGATGTTAAAGCACACATTAATTTTAAACATGSGGAAAGTTTAAAC 1997
          ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db      914  ATTAATAATATATATAAAAAAATTTAAAAATTTAAACATTTCTAAATTTCTTAAC 865
          ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

Search completed: August 19, 2003, 15:17:45
Job time : 581 secs

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